## SEQUENCE LISTING

Griffith, Irwin J Kuo, Mei-Chang Luqman, Mohammad

<12	0> ፐ	CEL	I. EP	<b>ΤΤΟΡ</b>	ES O	F RY	FGRA	99 D	ሰ፤.ፒ.ፑ	N 2AT.	LERĆ	FN.				
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							cag Gln									582
							gcc Ala									630
							ttc Phe 205									678
							aag Lys									726
gtc Val 230	aag Lys	cag Gln	gcc Ala	tac Tyr	gcc Ala 235	gcc Ala	acc Thr	gtc Val	gcc Ala	gcc Ala 240	gcg Ala	ccc Pro	gag Glu	gtc Val	aag Lys 245	774
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							ccc Pro									870
							gca Ala 285									918
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Ala Ala Glu Gly
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20

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Phe Ile Pro Ser
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Gln Ala Tyr Ala
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Ala Ala Ala Thr
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Thr Gly Ala Ala
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1 5 10 15

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                                                                    106
Val Gln Gln Tyr Thr Val Ala Leu Phe Leu Ala Val Ala Ser Cys Arg
gec ege gec tee tae gec gec gec gec gec tae gec eec gec act eec
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Ala Arg Ala Ser Tyr Ala Ala Asp Ala Gly Tyr Ala Pro Ala Thr Pro
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gcc acc ccg gct acc ccc gcg gcc cca ggc gca gcg gtg cca gca ggg
Ala Thr Pro Ala Thr Pro Ala Ala Pro Gly Ala Ala Val Pro Ala Gly
aag gcg gcg acc gag gag cag aag ctg atc gag aag atc aac gcc ggc
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ctg Leu 75	ggg Gly	gac Asp	ctc Leu	ccg Pro	acc. Thr 80	aac Asn	tac Tyr	gcc Ala	gat Asp	gtc Val 85	aac Asn	tcc Ser	agg Arg	gcc Ala	cag Gln 90	394
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aac Asn	aag L <b>y</b> s	gcc Ala	atc Ile 190	aag Lys	gag Glu	agc Ser	acg Thr	ggc Gly 195	ggc Gly	acc Thr	tac Tyr	gag Glu	agc Ser 200	tac Tyr	aag Lys	730
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gtc Val	gca Ala 220	Ser	gcg Ala	ccg Pro	gag Glu	gtc Val 225	aag Lys	tac Tyr	gcc Ala	gtc Val	ttt Phe 230	Glu	acc Thr	gcg Ala	ctg Leu	826
aaa Lys 235	aag Lys	gcg Ala	gtc Val	acc Thr	gcc Ala 240	Met	tcc Ser	gag Glu	gcc Ala	cag Gln 245	Lys	gaa Glu	gcc Ala	aag Lys	ccc Pro 250	874
gcc Ala	acc Thr	gcc Ala	acc Thr	ccg Pro 255	acc Thr	ccc Pro	acc Thr	gca Ala	act Thr 260	Ala	gcg Ala	gcc Ala	gcg Ala	gtg Val 265	Ala	922
acc Thr	aac Asn	gcc	gcc Ala 270	Pro	gtc Val	gct Ala	gct Ala	ggt Gly 275	ggc	tac Tyr	aaa Lys	atc :Ile	tga	tcaa	ctc	971

gctagcaata tacacatcca tcatgcacat atagagctgt gtatgtatgt gcatgcatgc 1031 cgtggcgccg cgcaagtttg ctcataatta attcttggtt ttcgttgctt gcatccacga 1091 gcgaccgagc ccgtggatag tcgcatgtgt atgtaattt ttctgagaaa tgtgtatatg 1151 taatatataa ttgagtacta aaaaaaaaaaa 1181

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Thr Pro Ala Thr Pro Ala Thr Pro Ala Ala Pro Gly Ala Ala Val Pro
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Ala Gly Phe Lys Ala Ala Val Ala Ala Ala Ala Gly Val Pro Pro Gly
45 50 55

Asp Lys Tyr Lys Thr Phe Val Glu Thr Phe Gly Lys Ala Ser Asn Lys
60 65 70

Ala Phe Leu Gly Asp Leu Pro Thr Asn Tyr Ala Asp Val Asn Ser Arg
75 80 85

Ala Gln Leu Thr Ser Lys Leu Asp Ala Ala Tyr Lys Leu Ala Tyr Asp 90 95 100

Ala Ala Gln Gly Ala Thr Pro Glu Ala Lys Tyr Asp Ala Tyr Val Ala 105 110 115 120

Thr Leu Ser Glu Ala Leu Arg Ile Ile Ala Gly Thr Leu Glu Val His 125 130 135

Ala Val Lys Pro Ala Ala Glu Glu Val Lys Pro Ile Pro Ala Gly Glu 140 145 150

Leu Gln Ile Val Asp Lys Ile Asp Val Ala Phe Arg Thr Ala Ala Thr 155 160 165

Ala Ala Asn Ala Ala Pro Thr Asn Asp Lys Phe Thr Val Phe Glu Thr 170 175 180

Thr Phe Asn Lys Ala Ile Lys Glu Ser Thr Gly Gly Thr Tyr Glu Ser 185 190 195 200

Tyr Lys Phe Ile Pro Thr Leu Glu Ala Ala Val Lys Gln Ala Tyr Ala 205 210 215 Ala Thr Val Ala Ser Ala Pro Glu Val Lys Tyr Ala Val Phe Glu Thr 220 225 230

Ala Leu Lys Lys Ala Val Thr Ala Met Ser Glu Ala Gln Lys Glu Ala 235 240 245

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